

## Class 11: Protein Alignments (2)

**Held:** Thursday, 1 October 2009

**Summary:** We continue our exploration of protein alignment by exploring some code that can be used for such alignment.

### Related Pages:

- EBoard.
- Due: Exploration of HIV Data.

### Notes:

- Readings for Tuesday: Chapter 5 and the HIV paper.
- One of the elementary schools in town was unexpectedly cancelled. We'll have children playing in the next room.

### Overview:

- Exploring Markham's data.
- Lab: Project 4.5.

## The HIV env. sequence

- We'd like to hear a short report from each group.
  - What question were you asking?
  - What data set (which sequences) did you choose for the exploration?
  - What did you learn or discover?
- We'd also like to hear about your experiences with alignment
  - What were the strengths and weaknesses of Clustal-W?
  - In what ways was it helpful that you'd written your own alignment program?
- What do you anticipate being the difficulties as you go from a program that aligns two sequences to one that aligns multiple sequences

## Lab

- Do Project 4.5.

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